RAW SEQUENCE LISTING PATENT APPLICATION US/08/977,221

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This Raw Listing contains the General Information Section and up to the first 5 pages.

1		SEQUENCE LISTING
2 3 4	(1) G	eneral Information:
5 6 7	(i)	APPLICANT: UNITED STATES OF AMERICA; DEPT OF HEALTH AND HUMAN SERVICES
8 9 10	(ii)	TITLE OF INVENTION: MOTILITY STIMULATING PROTEIN USEFUL IN CANCER DIAGNOSIS AND THERAPY
11 12 13	(iii)	NUMBER OF SEQUENCES: 69
14 15 16 17 18	(iv)	CORRESPONDENCE ADDRESS: (A) ADDRESSEE: MORGAN & FINNEGAN (B) STREET: 345 PARK AVENUE (C) CITY: NEW YORK (D) STATE: NEW YORK
19 20 21		(E) COUNTRY: U.S.A. (F) ZIP: 10154
22 23 24 25 26 27	(V)	COMPUTER READABLE FORM: (A) MEDIUM TYPE: Floppy Disk (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: WordPerfect 5.1
28 29 30 31 32	(vi)	CURRENT APPLICATION DATA: (A) APPLICATION NUMBER: (B) FILING DATE: (C) CLASSIFICATION:
33 34 35 36	(vii)	PRIOR APPLICATION DATA (A) APPLICATION NUMBER: 08/346,455 (B) FILING DATE: 28-NOV-1994
37 38 39 40	(vii)	PRIOR APPLICATION DATA (A) APPLICATION NUMBER: 08/249,182 (B) FILING DATE: 25-MAY-1994
41 42 43 44	(vii)	PRIOR APPLICATION DATA (A) APPLICATION NUMBER: 07/822,043 (B) FILING DATE: 17-JAN-1992
45 46	(viii)	ATTORNEY/AGENT INFORMATION: (A) NAME: DOROTHY R. AUTH

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REGISTRATION NUMBER: 36,434
47
              (B)
              (C) DOCKET NUMBER: 2026-4149US3
48
49
50
              TELECOMMUNICATION INFORMATION:
        (ix)
51
              (A) TELEPHONE: (212) 758-4800
52
              (B) TELEFAX: (212) 751-6849
53
54
55
56
57
58
59
60
61
    (2) INFORMATION FOR SEQ ID NO:1:
62
63
64
          (i) SEQUENCE CHARACTERISTICS:
65
               (A) LENGTH: 5
66
               (B) TYPE: amino acid
               (D) TOPOLOGY: linear
67
68
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
69
70
    Trp His Val Ala Arg
71
72
73
74
    (2) INFORMATION FOR SEQ ID NO:2:
75
          (i) SEQUENCE CHARACTERISTICS:
76
77
               (A) LENGTH: 6
               (B) TYPE: amino acid
78
               (D) TOPOLOGY: linear
79
80
81
82
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
83
84
    Pro Leu Asp Val Tyr Lys
      1
85
86
    (2) INFORMATION FOR SEQ ID NO:3:
87
88
          (i) SEQUENCE CHARACTERISTICS:
89
90
               (A) LENGTH: 5
91
               (B) TYPE: amino acid
92
               (D) TOPOLOGY: linear
93
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
94
95
96
    Tyr Pro Ala Phe Lys
97
      1
98
99
    (2) INFORMATION FOR SEQ ID NO:4:
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100
           (i) SEQUENCE CHARACTERISTICS:
101
102
                (A) LENGTH: 5 amino acids
                (B) TYPE: amino acid
103
                (D) TOPOLOGY: linear
104
105
106
107
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
108
     Gln Ala Glu Val Ser
109
      1
110
111
     (2) INFORMATION FOR SEQ ID NO:5:
112
113
114
           (i) SEQUENCE CHARACTERISTICS:
                (A) LENGTH: 10
115
                (B) TYPE: amino acid
116
117
118
119
120
121
122
123
124
                (D) TOPOLOGY: linear
125
126
127
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
128
129
130
     Pro Glu Glu Val Thr Arg Pro Asn Tyr Leu
131
       1
                        5
132
133
     (2) INFORMATION FOR SEQ ID NO:6:
134
           (i) SEQUENCE CHARACTERISTICS:
135
                (A) LENGTH: 9
136
                (B) TYPE: amino acid
137
138
                (D) TOPOLOGY: linear
139
140
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
141
     Tyr Asp Val Pro Trp Asn Glu Thr Ile
142
143
       1
                        5
144
     (2) INFORMATION FOR SEQ ID NO:7:
145
146
147
          (i) SEQUENCE CHARACTERISTICS:
148
                (A) LENGTH: 10
149
                (B) TYPE: amino acid
                (D) TOPOLOGY: linear
150
151
152
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153
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
154
     Val Pro Pro Phe Glu Asn Ile Glu Leu Tyr
155
156
157
      (2) INFORMATION FOR SEQ ID NO:8:
158
159
           (i) SEQUENCE CHARACTERISTICS:
160
161
                (A) LENGTH: 11
                (B) TYPE: amino acid
162
163
                (D) TOPOLOGY: linear
164
165
166
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
167
168
     Gly Gly Gln Pro Leu Trp Ile Thr Ala Thr Lys
169
170
     (2) INFORMATION FOR SEQ ID NO:9:
171
172
173
           (i) SEQUENCE CHARACTERISTICS:
174
                (A) LENGTH: 16
                (B) TYPE: amino acid
175
                (D) TOPOLOGY: linear
176
177
178
179
180
181
182
183
184
185
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
186
     Val Asn Ser Met Gln Thr Val Phe Val Gly Tyr Gly
187
188
       1
                        5
                                             10
189
190
     Pro Thr Phe Lys
191
               15
192
      (2) INFORMATION FOR SEQ ID NO:10:
193
194
           (i) SEQUENCE CHARACTERISTICS:
195
196
                (A) LENGTH: 12
                (B) TYPE: amino acid
197
198
                (D) TOPOLOGY: linear
199
200
201
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:
202
203
     Asp Ile Glu His Leu Thr Ser Leu Asp Phe Phe Arg
204
                        5
205
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206
     (2) INFORMATION FOR SEQ ID NO:11:
207
208
           (i) SEQUENCE CHARACTERISTICS:
                (A) LENGTH: 23
209
210
                (B) TYPE: amino acid
                (D) TOPOLOGY: linear
211
212
213
214
215
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:
216
217
     Thr Glu Phe Leu Ser Asn Tyr Leu Thr Asn Val Asp
218
219
     Asp Ile Thr Leu Val Pro Glu Thr Leu Gly Arg
220
221
222
223
     (2) INFORMATION FOR SEQ ID NO:12:
224
225
226
           (i) SEQUENCE CHARACTERISTICS:
227
                (A) LENGTH: 18
                (B) TYPE: nucleic acid
228
                (C) STRANDEDNESS: single
229
230
                (D) TOPOLOGY: linear
231
232
233
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:
234
235
     GTTGGCAGCN ACRTGCCA
                                                               18
236
     (2) INFORMATION FOR SEQ ID NO:13:
237
238
239
240
241
242
243
244
245
246
           (i) SEQUENCE CHARACTERISTICS:
247
                (A) LENGTH: 18
248
                (B) TYPE: nucleic acid
                (C) STRANDEDNESS: single
249
250
                (D) TOPOLOGY: linear
251
252
253
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:
254
                                                               18
255
     TGGCAYGTNG CTGCCAAC
256
257
     (2) INFORMATION FOR SEQ ID NO:14:
258
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SEQUENCE VERIFICATION REPORT PATENT APPLICATION *US/08/977,221*

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